



1/6

SEQUENCE LISTING

RECEIVED
DEC 13 2001
TECH CENTER 1600/2300

<110> James M. Anderson
Christina M. Van Itallie

<120> Human Occludin, Its Uses and Enhancement of Drug
Absorption Using Occludin Inhibitors

<130> OCR-754.CIP

<140> US 09/891,064

<141> 2001-06-25

<150> US 09/142,732

<151> 1998-09-15

<160> 6

<170> MS DOS

<210> 1

<211> 2312

<212> DNA

<213> Homo sapiens

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<221> mat_peptide

<222> complete sequence

<223> human occludin

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<210>      2
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<212>      PRT
<213>      Homo sapiens
<220>
<221>      peptide
<222>      complete sequence
<223>      human occludin
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Glu Phe Lys Pro Asn His Tyr Ala Pro Ser Asn Asp Ile Tyr Gly
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Gly Glu Met His Val Arg Pro Met Leu Ser Gln Pro Ala Tyr Ser
      35                      40                      45

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Phe Tyr Pro Glu Asp Glu Ile Leu His	Phe Tyr Lys Trp Thr Ser
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Pro Pro Gly Val Ile Arg Ile Leu Ser	Met Leu Ile Ile Val Met
65	70 75
Cys Ile Ala Ile Phe Ala Cys Val Ala	Ser Thr Leu Ala Trp Asp
80	85 90
Arg Gly Tyr Gly Thr Ser Leu Leu Gly	Gly Ser Val Gly Tyr Pro
95	100 105
Tyr Gly Gly Ser Gly Phe Gly Ser Tyr	Gly Ser Gly Tyr Gly Tyr
110	115 120
Gly Tyr Gly Tyr Gly Tyr Gly Tyr Gly	Gly Tyr Thr Asp Pro Arg
125	130 135
Ala Ala Lys Gly Phe Met Leu Ala Met	Ala Ala Phe Cys Phe Ile
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Ala Ala Leu Val Ile Phe Val Thr Ser	Val Ile Arg Ser Glu Met
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Ser Arg Thr Arg Arg Tyr Tyr Leu Ser	Val Ile Ile Val Ser Ala
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Ile Leu Gly Ile Met Val Phe Ile Ala	Thr Ile Val Tyr Ile Met
185	190 195
Gly Val Asn Pro Thr Ala Gln Ser Ser	Gly Ser Leu Tyr Gly Ser
200	205 210
Gln Ile Tyr Ala Leu Cys Asn Gln Phe	Tyr Thr Pro Ala Ala Thr
215	220 225
Gly Leu Tyr Val Asp Gln Tyr Leu Tyr	His Tyr Cys Val Val Asp
230	235 240
Pro Gln Glu Ala Ile Ala Ile Val Leu	Gly Phe Met Ile Ile Val
245	250 255
Ala Phe Ala Leu Ile Ile Phe Phe Ala	Val Lys Thr Arg Arg Lys
260	265 270
Met Asp Arg Tyr Asp Lys Ser Asn Ile	Leu Trp Asp Lys Glu His
275	280 285
Ile Tyr Asp Glu Gln Pro Pro Asn Val	Glu Glu Trp Val Lys Asn
290	295 300
Val Ser Ala Gly Thr Gln Asp Val Pro	Ser Pro Pro Ser Asp Tyr
305	310 315
Val Glu Arg Val Asp Ser Pro Met Ala	Tyr Ser Ser Asn Gly Lys
320	325 330

Val Asn Asp Lys Arg Phe Tyr Pro Glu Ser Ser Tyr Lys Ser Thr		
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Pro Val Pro Glu Val Val Gln Glu Leu Pro Leu Thr Ser Pro Val		
	350	355 360
Asp Asp Phe Arg Gln Pro Arg Tyr Ser Ser Gly Gly Asn Phe Glu		
	365	370 375
Thr Pro Ser Lys Arg Ala Pro Ala Lys Gly Arg Ala Gly Arg Ser		
	380	385 390
Lys Arg Thr Glu Gln Asp His Tyr Glu Thr Asp Tyr Thr Thr Gly		
	395	400 405
Gly Glu Ser Cys Asp Glu Leu Glu Glu Asp Trp Ile Arg Glu Tyr		
	410	415 420
Pro Pro Ile Thr Ser Asp Gln Gln Arg Gln Leu Tyr Lys Arg Asn		
	425	430 435
Phe Asp Thr Gly Leu Gln Glu Tyr Lys Ser Leu Gln Ser Glu Leu		
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Asp Glu Ile Asn Lys Glu Leu Ser Arg Leu Asp Lys Glu Leu Asp		
	455	460 465
Asp Tyr Arg Glu Glu Ser Glu Glu Tyr Met Ala Ala Ala Asp Glu		
	470	475 480
Tyr Asn Arg Leu Lys Gln Val Lys Gly Ser Ala Asp Tyr Lys Ser		
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Lys Lys Asn His Cys Lys Gln Leu Lys Ser Lys Leu Ser His Ile		
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Lys Lys Met Val Gly Asp Tyr Asp Arg Gln Lys Thr		
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<210> 3

<211> 24

<212> PRT

<213> Artificial Sequence

<220>

<221> peptide

<223> construct used in experiments

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5/6

Cys Asp Arg Gly Tyr Gly Thr Ser Leu Leu Gly Gly Ser Val Gly
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Tyr Pro Tyr Gly Gly Ser Gly Phe Gly
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<210> 4

<211> 24

<212> PRT

<213> Artificial Sequence

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<223> construct used in experiments

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Cys Ser Tyr Gly Ser Gly Tyr Gly Tyr Gly Tyr Gly Tyr Gly Tyr
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Gly Tyr Gly Gly Tyr Thr Asp Pro Arg
20

<210> 5

<211> 20

<212> PRT

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Asn His Tyr Ala Pro Ser Asn Asp Ile Tyr Gly Gly Glu Met Val
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His Arg Pro Met Leu
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<210> 6

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<212> PRT

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Ala Ser Gln Gln Val Tyr Arg Lys Asp Pro Cys
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